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By: [Signature]
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Docket No. PE 0195-1 DIV

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IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

Applicants: Jennifer L. Hillman, Surya K. Goli
Title: A NOVEL PROSTATE-ASSOCIATED KALLIKREIN
Serial No.: 09/170,980 Filed: Herewith
Examiner: Y. Eyler Group Art Unit: 1642

11/C
M. Q.
10/30/00

Box Non-Fee Amendment
Commissioner for Patents
Washington, D.C. 20231

SUBSTITUTE SUBMISSION UNDER 37 CFR §1.821- 1.825 SEQUENCE LISTING

Sir:

This communication is in response to the Notice to Comply mailed by the U.S. Patent and Trademark Office on September 8, 2000. A copy of the Notice to Comply accompanies this statement.

In accordance with the requirements of 37 CFR § 1.821-1.825, Applicants hereby submit one (1) substitute diskette containing the computer-readable information for the Substitute Sequence Listing of the above-identified application. The diskette complies with the requirements of 37 CFR § 1.824 and is IBM PC compatible using a UNIX operating system with PERL Program.

Enclosed is a paper copy of the Substitute Sequence Listing.

The content of the Substitute Sequence Listing paper copy is identical to the computer-readable copy, as required under 37 CFR § 1.821(f).

Respectfully submitted,

INCYTE GENOMICS, INC.

Date:

10 October 2000

[Signature]

Diana Hamlet-Cox, Ph.D.

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SEQUENCE LISTING



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<110> Hillman, Jennifer L.
Goli, Surya K.

<120> A NOVEL PROSTATE-ASSOCIATED KALLIKREIN

<130> PF-0195-1 DIV

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<141> 1998-10-13

<150> 08/790,137

<151> 1997-01-29

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50 55 60
Leu Thr Ala Ala His Cys Leu Lys Pro Arg Tyr Ile Val His Leu
65 70 75
Gly Gln His Asn Leu Gln Lys Glu Glu Gly Cys Glu Gln Thr Arg
80 85 90
Thr Ala Thr Glu Ser Phe Pro His Pro Gly Phe Asn Asn Ser Leu
95 100 105
Pro Asn Lys Asp His Arg Asn Asp Ile Met Leu Val Lys Met Ala
110 115 120
Ser Pro Val Ser Ile Thr Trp Ala Val Arg Pro Leu Thr Leu Ser
125 130 135
Ser Arg Cys Val Thr Ala Gly Thr Ser Cys Leu Ile Ser Gly Trp
140 145 150
Gly Ser Thr Ser Ser Pro Gln Leu Arg Leu Pro His Thr Leu Arg
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Cys Ala Asn Ile Thr Ile Ile Glu His Gln Lys Cys Glu Asn Ala
170 175 180
Tyr Pro Gly Asn Ile Thr Asp Thr Met Val Cys Ala Ser Val Gln
185 190 195
Glu Gly Gly Lys Asp Ser Cys Gln Gly Asp Ser Gly Gly Pro Leu
200 205 210
Val Cys Asn Gln Ser Leu Gln Gly Ile Ile Ser Trp Gly Gln Asp
215 220 225
Pro Cys Ala Ile Thr Arg Lys Pro Gly Val Tyr Thr Lys Val Cys
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Lys Tyr Val Asp Trp Ile Gln Glu Thr Met Lys Asn Asn
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cgacgntcat ngcccccaga tggttcctga cagcagccca ctgcctnaag ccccgctaca 240
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Leu Thr Ala Ala His Cys Ile Ser Asp Asn Tyr Gln Leu Trp Leu
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Gly Arg His Asn Leu Phe Asp Asp Glu Asn Thr Ala Gln Phe Val
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His Val Ser Glu Ser Phe Pro His Pro Gly Phe Asn Met Ser Leu
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Leu Glu Asn His Thr Arg Gln Ala Asp Glu Asp Tyr Ser His Asp
110 115 120
Leu Met Leu Leu Arg Leu Thr Glu Pro Ala Asp Thr Ile Thr Asp
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Ala Val Lys Val Val Glu Leu Pro Thr Gln Glu Pro Glu Val Gly
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Ser Thr Cys Leu Ala Ser Gly Trp Gly Ser Ile Glu Pro Glu Asn
155 160 165

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Phe	Met	Leu	Cys	Val	Gly	His	Leu	Glu	Gly	Gly	Lys	Asp	Thr	Cys
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Val	Gly	Asp	Ser	Gly	Gly	Pro	Leu	Met	Cys	Asp	Gly	Val	Leu	Gln
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Gly	Val	Thr	Ser	Trp	Gly	Tyr	Val	Pro	Cys	Gly	Thr	Pro	Asn	Lys
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Pro	Ser	Val	Ala	Val	Arg	Val	Leu	Ser	Tyr	Val	Lys	Trp	Ile	Glu
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Cys	Val	Gly	Asp	Ser	Gly	Gly	Pro	Leu	Ile	Cys	Asp	Gly	Val	Leu
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Gln	Gly	Ile	Thr	Ser	Trp	Gly	Pro	Thr	Pro	Cys	Ala	Leu	Pro	Asn
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Val	Pro	Gly	Ile	Tyr	Thr	Lys	Leu	Ile	Glu	Tyr	Arg	Ser	Trp	Ile
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				260										

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